Molecular epidemiology of simian malaria causing zoonosis in Vietnam

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Concerted control measures have considerably reduced the burden of malaria in Vietnam, and the parasites that cause it are now mostly restricted to forested rural areas. The high risk of malaria infection associated with subsistence activity in forests is a well-known and widely reported phenomenon in Southeast Asia. This phenomenon is commonly referred to as 'forest malaria'. The transmission of P. knowlesi from monkeys to humans contributes an additional complication to our understanding of forest malaria. However, zoonotic malaria by P. knowlesi infection in those areas was unclear. In order to determine whether P. knowlesi is infecting the human population of the Khanh Phu Commune, Khanh Vinh District, Khanh Hoa Province, we collected mosquito and human blood samples from the commune, and analyzed these for the presence of malaria parasites using PCR. In analysis of sporozoite positive salivary glands, P. knowlesi was detected in 42 salivary glands, which was the same detection number as P. falciparum in the samples. Mixed infection with P. knowlesi and human malaria parasites was detected in 27 samples, 19 were mixed with P. vivax, one with P. falciparum, and 7 with P. vivax and P. falciparum. The results suggest that Anopheles dirus harbors P. knowlesi in a considerably high rate, and bites monkeys as frequently as humans. In the analysis of human blood samples, P. knowlesi was detected in 45 out of 170 blood samples with malaria parasites. All P. knowlesi parasites were found in mixed infections with other human malaria parasites. Gametocytes of P. falciparum and P. vivax, moreover, were detected by RT-PCR analysis but that of P. knowlesi was not detected. These results indicate that natural human infection of P. knowlesi is prevalent in this area. In this study, we were not able to determine whether P. knowlesi is being transmitted from human to human although transmission from monkey to humans via mosquitoes seems highly likely.